

2/33 ᇤ 26S mRNA Flips back into cytoplasm? Initiates E1 translocation Internal cleavable signal No additional function known Generates 6K and E1 C-terminus of 6K The E1 signal AAA Membrane protein E1 Flips back into cytoplasm? Initiates 6K translocation Internal cleavable signal Generates p62 and 6K Nucleocapsid binding 6K C-terminus of p62 The 6K signal Membrane protein 6K Membrane protein p62 Initiates p62 translocation after cleavage of capsid Uncleaved (translocates into lumen) p62-E1 oligomerization Becomes glycosylated N-terminus of p62 The p62 signal প্ত p62 Capsid protein CAPSID m7GI

Figure 2

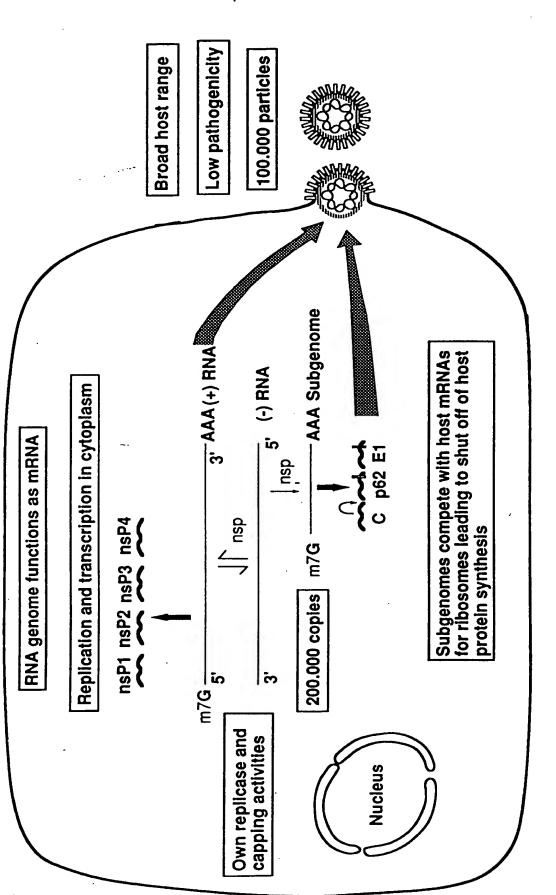


Figure 3

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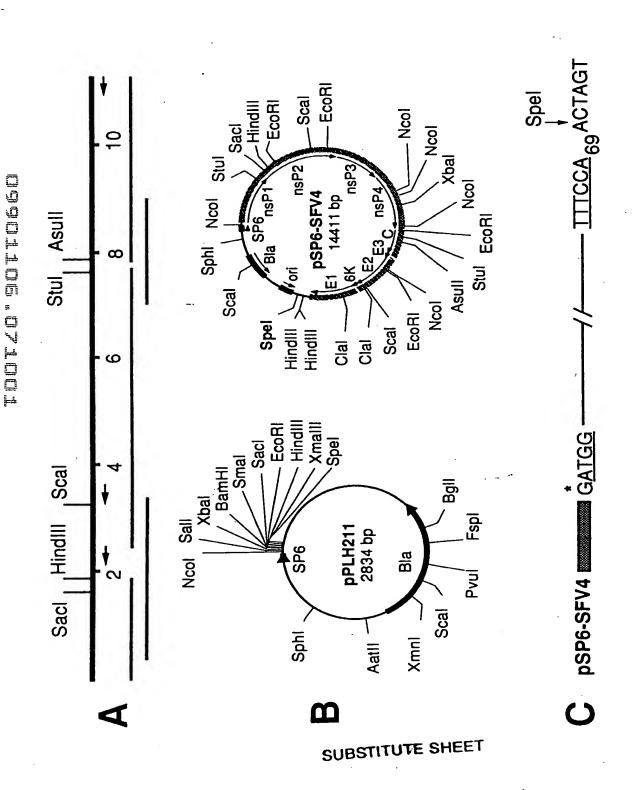


Figure 4

175

Figure 5 (1)

GCTACGCGAG AGATTAACCA CCCACG ATG GCC GCC AAA GTG CAT GTT GAT ATT Met Ala Ala Lys Val His Val Asp Ile GAG GCT GAC AGC CCA TTC ATC AAG TCT TTG CAG AAG GCA TTT CCG 158 Glu Ala Asp Ser Pro Phe Ile Lys Ser Leu Gln Lys Ala Phe Pro 15 TCG TTC GAG GTG GAG TCA TTG CAG GTC ACA CCA AAT GAC CAT GCA Ser Phe Glu Val Glu Ser Leu Gln Val Thr Pro Asn Asp His Ala 30 AAT GCC AGA GCA TTT TCG CAC CTG GCT ACC AAA TTG ATC GAG CAG Asn Ala Arg Ala Phe Ser His Leu Ala Thr Lys Leu Ile Glu Gln GAG ACT GAC AAA GAC ACA CTC ATC TTG GAT ATC GGC AGT GCG CCT Glu Thr Asp Lys Asp Thr Leu Ile Leu Asp Ile Gly Ser Ala Pro TCC AGG AGA ATG ATG TCT ACG CAC AAA TAC CAC TGC GTA TGC CCT 338 Ser Arg Arg Met Met Ser Thr His Lys Tyr His Cys Val Cys Pro 75 ATG CGC AGC GCA GAA GAC CCC GAA AGG CTC GAT AGC TAC GCA AAG Met Arg Ser Ala Glu Asp Pro Glu Arg Leu Asp Ser Tyr Ala Lys 85 90 AAA CTG GCA GCG GCC TCC GGG AAG GTG CTG GAT AGA GAG ATC GCA Lys Leu Ala Ala Ala Ser Gly Lys Val Leu Asp Arg Glu Ile Ala 100 105 GGA AAA ATC ACC GAC CTG CAG ACC GTC ATG GCT ACG CCA GAC GCT Gly Lys Ile Thr Asp Leu Gln Thr Val Met Ala Thr Pro Asp Ala 115 120 GAA TCT CCT ACC TTT TGC CTG CAT ACA GAC GTC ACG TGT CGT ACG Glu Ser Pro Thr Phe Cys Leu His Thr Asp Val Thr Cys Arg Thr 130 135 GCA GCC GAA GTG GCC GTA TAC CAG GAC GTG TAT GCT GTA CAT GCA 563 Ala Ala Glu Val Ala Val Tyr Gln Asp Val Tyr Ala Val His Ala 155 150 CCA ACA TCG CTG TAC CAT CAG GCG ATG AAA GGT GTC AGA ACG GCG Pro Thr Ser Leu Tyr His Gln Ala Met Lys Gly Val Arg Thr Ala 165 TAT TGG ATT GGG TTT GAC ACC CCG TTT ATG TTT GAC GCG CTA Tyr Trp Ile Gly Phe Asp Thr Thr Pro Phe Met Phe Asp Ala Leu

GATGGCGGAT GTGTGACATA CACGACGCCA AAAGATTTTG TTCCAGCTCC TGCCACCTCC

180

185

Gly		ACC Thr 195	Tyr			Ala		CAG Gln	698
Leu		AAC Asn 210	Ile			Ala			743
Gly		AAA Lys 225	Leu			Lys			788
Pro		GTC Val 240	Met			Ser			833
Glu		CTA Leu 255				Leu			878
His		AAA Lys 270				Arg			923
Val		GGG Gly 285							968
Gly		AAA Lys 300							1013
Glu		GTG Val 315							1058
		CCT Pro 330							1103
		GGC Gly 345						GAG Glu	1148
		TTA Leu 360							1193
		CGA Arg 375							1238
		GTC Val 390							1283

Figure 5 (3)

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			CTT												1328
	Ala	Asp	Leu	Asp		Glu	Lys	Pro	Leu		Val	Arg	Glu	Arg	
400					405					410					
TCA	СТТ	АСТ	TGC	TGC	TGC	TTG	TGG	GCA	TTT	AAA	ACG	AGG	AAG	ATG	1373
Ser	Leu	Thr	Cys	Cys	Cys	Leu	Trp	Ala	Phe	Lys	Thr	Arg	Lys	Met	
415					420					425					
							~~~	100	C3.C	202	N CON	CITY:	N N C	CTC	1418
CAC	ACC	ATG	TAC Tyr	AAG	AAA	Pro	CAC	Thr	Gln	Thr	Ile	Val	Lvs	Val	1410
430	1111	Mec	TYL	Ly 3	435		пор			440					
			TTT												1463
_	Ser	Glu	Phe	Asn		Phe	Val	He	Pro	Ser 455	Leu	Trp	Ser	Thr	
445					450					433					
GGC	CTC	GCA	ATC	CCA	GTC	AGA	TCA	CGC	ATT	AAG	ATG	CTT	TTG	GCC	1508
			Ile												
460					465					470					
											~		<b>m</b> 00	ma.	1550
														TCA	1553
Lys 475	_	Thr	Lys	Arg	480	Leu	Tre	Pro	vai	485	Asp	AIG	Ser	361	
413					400					•••					
														GAG	1598
Ala	Arg	Asp	Ala	Glu	Gln	Glu	Glu	Lys	Glu	Arg	Leu	Glu	Ala	Glu	
490					495					500					-
omo	<b>3.</b> Cm	202	C	ccc	מידיא	CCA	ccc	CTC	CTC	CCC	ልጥና	GCG	CCG	GCG	1643
			GAA												2010
505		ALG	GIU	AIG	510					515					
														CAC	1688
Glu	Thr	Gly	Val	Val	_	Val	Asp	Val	Glu		Leu	Glu	Tyr	His	
520					525					530					
CCA	ርርጥ	GC A	GGG	GTC	GTG	GAA	ACA	ССТ	CGC	AGC	GCG	TTG	AAA	GTC	1733
			Gly												
535	_		•		540					545					
															1770
														CTG	1778
		Gln	Pro	Asn	<b>Asp</b> 555	Val	Leu	Leu	GIY	560	туг	vaı	vai	Leu	
550					333					500					
														CAC	1823
Ser	Pro	Gln	Thr	Val		Lys	Ser	Ser	Lys		Ala	Pro	Val	His	
565					570					575					
ር ር ጥ	ርጥአ	GC N	CAC	CAG	GTG	AAA	АТА	ATA	ACA	CAT	AAC	GGG	AGG	GCC	1868
														Ala	
580					585	-				590					
										<b>i</b>	ama	~~ <u>`</u>	~m>	CC3	1017
GGC	GGT	TAC	CAG	GTC	GAC	GGA	TAT	GAC	GGC	AGG	GIC V=1	CTA t.e.u	CTA Leu	CCA	1913
_	_	чуr	GIn	val	Asp 600	GIÀ	ıyr	изр	GIA	605	val	Leu	Dea	Pro	
595							~-	ioe1			SHE	ΞT			

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# Figure 5 (4)

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	Gly				CCG Pro 615									GAG Glu	1958
	Ala				TAC Tyr 630									AAA Lys	2003
	Tyr				GTT Val 645									GAG Glu	2048
	Asn				GTC Val 660									TAC Tyr	2093
	Phe				AAA Lys 675									GCG Ala	2138
					GTG Val 690										2183
Glu 700	Phe	Ala	Tyr	Glu	Gly 705	Leu	Lys	Ile	Arg	Pro 710	Ser	Ala	Pro	_	2228
Lys 715	Thr	Thr	Val	Val	Gly 720	Val	Phe	Gly	Val	Pro 725	Gly	Ser	Gly	_	2273
Ser 730	Ala	Ile	Ile	Lys	AGC Ser 735	Leu	Val	Thr	Lys	His 740	Asp	Leu	Val	Thr	2318
Ser 745	Gly	Lys	Lys	Glu	AAC Asn 750	Cys	Gln	Glu	Ile	Val 755	Asn	Asp	Val	Lys	2363
Lys 760	His	Arg	Gly	Lys	Gly 765	Thr	Ser	Arg	Glu	Asn 770	Ser	Asp	Ser		2408
					CGT Arg 780										2453
					CAT His 795									GCT Ala	2498
					AGC Ser 810										2543

### CAA TGC GGA TTC TTC AAT ATG ATG CAG CTT AAG GTG AAC TTC AAC Gln Cys Gly Phe Phe Asn Met Met Gln Leu Lys Val Asn Phe Asn 825 830 CAC AAC ATC TGC ACT GAA GTA TGT CAT AAA AGT ATA TCC AGA CGT His Asn Ile Cys Thr Glu Val Cys His Lys Ser Ile Ser Arg Arg TGC ACG CGT CCA GTC ACG GCC ATC GTG TCT ACG TTG CAC TAC GGA 2678 Cys Thr Arg Pro Val Thr Ala Ile Val Ser Thr Leu His Tyr Gly 850 855 GGC AAG ATG CGC ACG ACC AAC CCG TGC AAC AAA CCC ATA ATC ATA Gly Lys Met Arg Thr Thr Asn Pro Cys Asn Lys Pro Ile Ile Ile 865 870 875 GAC ACC ACA GGA CAG ACC AAG CCC AAG CCA GGA GAC ATC GTG TTA Asp Thr Thr Gly Gln Thr Lys Pro Lys Pro Gly Asp Ile Val Leu 880 ACA TGC TTC CGA GGC TGG GCA AAG CAG CTG CAG TTG GAC TAC CGT Thr Cys Phe Arg Gly Trp Ala Lys Gln Leu Gln Leu Asp Tyr Arg 895 900 GGA CAC GAA GTC ATG ACA GCA GCA GCA TCT CAG GGC CTC ACC CGC Gly His Glu Val Met Thr Ala Ala Ala Ser Gln Gly Leu Thr Arg 915 AAA GGG GTA TAC GCC GTA AGG CAG AAG GTG AAT GAA AAT CCC TTG 2903 Lys Gly Val Tyr Ala Val Arg Gln Lys Val Asn Glu Asn Pro Leu 930 TAT GCC CCT GCG TCG GAG CAC GTG AAT GTA CTG CTG ACG CGC ACT Tyr Ala Pro Ala Ser Glu His Val Asn Val Leu Leu Thr Arg Thr 945 GAG GAT AGG CTG GTG TGG AAA ACG CTG GCC GGC GAT CCC TGG ATT Glu Asp Arg Leu Val Trp Lys Thr Leu Ala Gly Asp Pro Trp Ile 960 965 AAG GTC CTA TCA AAC ATT CCA CAG GGT AAC TTT ACG GCC ACA TTG 3038 Lys Val Leu Ser Asn Ile Pro Gln Gly Asn Phe Thr Ala Thr Leu 970 975 GAA GAA TGG CAA GAA GAA CAC GAC AAA ATA ATG AAG GTG ATT GAA Glu Glu Trp Gln Glu Glu His Asp Lys Ile Met Lys Val Ile Glu 985 990 GGA CCG GCT GCG CCT GTG GAC GCG TTC CAG AAC AAA GCG AAC GTG Gly Pro Ala Ala Pro Val Asp Ala Phe Gln Asn Lys Ala Asn Val 1000 1005 TGT TGG GCG AAA AGC CTG GTG CCT GTC CTG GAC ACT GCC GGA ATC Cys Trp Ala Lys Ser Leu Val Pro Val Leu Asp Thr Ala Gly Ile 1015 1020

### Figure 5 (6)

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Leu		A GAG a Glu I	Trp			Thr		
Asp		TAC Tyr				a Asn		326:
Lys		GGA Gly				<b>Leu</b>		3308
Lys		CTG Leu				Trp		3353
Gly		ATG Met				Thr		3398
Glu		CAT His				Trp		3443
Gln		ATC Ile-				Pro		3488
Asp		ATT Ile 1				Pro		3533
Ala		AAG Lys 1				Val		3578
Asn		AGA Arg				Val		3623
		CCT Pro			Thr			3668
		GCC (Ala 2			Asp			3713
		GGC 2 Gly 2			Val			3758
lu		ATC (			Gln			3803

1435

#### ATG AAG CTG CAG ATG CTT GGG GGA GAT GCG CTA CGA CTG CTA AAA Met Lys Leu Gln Met Leu Gly Gly Asp Ala Leu Arg Leu Lys 1245 1250 CCC GGC GGC ATC TTG ATG AGA GCT TAC GGA TAC GCC GAT AAA ATC Pro Gly Gly Ile Leu Met Arg Ala Tyr Gly Tyr Ala Asp Lys Ile 1260 1265 AGC GAA GCC GTT GTT TCC TCC TTA AGC AGA AAG TTC TCG TCT GCA Ser Glu Ala Val Val Ser Ser Leu Ser Arg Lys Phe Ser Ser Ala 1270 1275 1280 AGA GTG TTG CGC CCG GAT TGT GTC ACC AGC AAT ACA GAA GTG TTC Arg Val Leu Arg Pro Asp Cys Val Thr Ser Asn Thr Glu Val Phe 1285 1290 1295 TTG CTG TTC TCC AAC TTT GAC AAC GGA AAG AGA CCC TCT ACG CTA Leu Leu Phe Ser Asn Phe Asp Asn Gly Lys Arg Pro Ser Thr Leu 1300 1305 CAC CAG ATG AAT ACC AAG CTG AGT GCC GTG TAT GCC GGA GAA GCC His Gln Met Asn Thr Lys Leu Ser Ala Val Tyr Ala Gly Glu Ala 1315 1320 ATG CAC ACG GCC GGG TGT GCA CCA TCC TAC AGA GTT AAG AGA GCA Met His Thr Ala Gly Cys Ala Pro Ser Tyr Arg Val Lys Arg Ala 1335 GAC ATA GCC ACG TGC ACA GAA GCG GCT GTG GTT AAC GCA GCT AAC Asp Ile Ala Thr Cys Thr Glu Ala Ala Val Val Asn Ala Ala Asn 1350 GCC CGT GGA ACT GTA GGG GAT GGC GTA TGC AGG GCC GTG GCG AAG Ala Arg Gly Thr Val Gly Asp Gly Val Cys Arg Ala Val Ala Lys 1360 1365 1370 AAA TGG CCG TCA GCC TTT AAG GGA GCA GCA ACA CCA GTG GGC ACA Lys Trp Pro Ser Ala Phe Lys Gly Ala Ala Thr Pro Val Gly Thr 1375 1380 ATT AAA ACA GTC ATG TGC GGC TCG TAC CCC GTC ATC CAC GCT GTA Ile Lys Thr Val Met Cys Gly Ser Tyr Pro Val Ile His Ala Val 1390 1395 1400 GCG CCT AAT TTC TCT GCC ACG ACT GAA GCG GAA GGG GAC CGC GAA Ala Pro Asn Phe Ser Ala Thr Thr Glu Ala Glu Gly Asp Arg Glu 1405 1410 TTG GCC GCT GTC TAC CGG GCA GTG GCC GCC GAA GTA AAC AGA CTG Leu Ala Ala Val Tyr Arg Ala Val Ala Ala Glu Val Asn Arg Leu 1420 1425 1430

#### SUBSTITUTE SHEET

1445

TCA CTG AGC AGC GTA GCC ATC CCG CTG CTG TCC ACA GGA GTG TTC Ser Leu Ser Ser Val Ala Ile Pro Leu Leu Ser Thr Gly Val Phe

1440

### Figure 5 (8)

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AGC GGC GGA AGA GAT AGG CTG CAG CAA TCC CTC AAC CAT CTA TTC Ser Gly Gly Arg Asp Arg Leu Gln Gln Ser Leu Asn His Leu Phe 1450  ACA GCA ATG GAC GCC ACG GAC GCT GAC GTG ACC ATC TAC TGC AGA Thr Ala Met Asp Ala Thr Asp Ala ASP Val Thr Ile Tyr Cys Arg 1465  ACA GCA AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG ASp Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480  ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1495  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT GGT GGT GTG Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG GCA GTG CCC TGC CTG TGC 4883
1450  ACA GCA ATG GAC GCC ACG GAC GCT GAC GTG ACC ATC TAC TGC AGA 4523 Thr Ala Met Asp Ala Thr Asp Ala Asp Val Thr Ile Tyr Cys Arg 1465  GAC AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG 4568 Asp Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480  ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC 4613 Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1495  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658 Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1520  TAC AGT ACC ACT GAC GGG TCG CTG TAC TGC TAC TTT GAA GGT ACG 4703 Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1535  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 4748 Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAA GAA CAA ATA CTG ACG TTG 4793 Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838 Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC CCC AGG ACA GTG CCC TGC CTG CTG CTG TGC 4883
ACA GCA ATG GAC GCC ACG GAC GCT GAC GTG ACC ATC TAC TGC AGA Thr Ala Met Asp Ala Thr Asp Ala Asp Val Thr Ile Tyr Cys Arg 1465  GAC AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG Asp Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480  ACG GCT GTG GAG TTG CTC AAT GAC GTG GAG CTG ACC ACA GAC Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1495  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAA GAA CAA ATA CTG ACG TTG Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC CCC AGG ACA GTG CCC TGC CTG TGC CAB8  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC CAB8  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG CTG TGC  A850  AST TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG CTG TGC  A850  A523  ATT TAC TAC TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC  A561  A571  A572  A573  A574  A575  A574  A57
Thr Ala Met Asp Ala Thr Asp Ala Asp Val Thr Ile Tyr Cys Arg 1475  GAC AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG Asp Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480  ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC ATT Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1500  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT CAT AAG GGC Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG ACG TYr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1530  AAA TTC AAC AGG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAA ATA GAA TGT CG GTG AAC 4838  CTG GGC GAA ACA ATG GAC AAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TTG CTG CAG ACA GAG ATA CTG AAC 4838
Thr Ala Met Asp Ala Thr Asp Ala Asp Val Thr Ile Tyr Cys Arg 1475  GAC AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG Asp Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480  ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC ATT Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1500  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT CAT AAG GGC Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG ACG TYr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1530  AAA TTC AAC AGG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAA ATA GAA TGT CG GTG AAC 4838  CTG GGC GAA ACA ATG GAC AAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TTG CTG CAG ACA GAG ATA CTG AAC 4838
1465       1470       1475         GAC AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG ASP Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480       4568         ASP Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480       1485       1490         ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC ATT Thr Ala Val Glu Leu Leu Asp Asp Asp Val Glu Leu Thr Thr Asp 1495       4613         TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT GGT CGT AAG GGC Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510       4658         TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG TYR Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1530       4703         TAC AGT AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asp Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540       1545         TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG TTP ATG Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555       1560         TCG CGC GAA ACA ATG GAC AAC AAC GAA CAG ATC CAAA TGT CCG GTG AAC 4838       4838         Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570       1757         GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
GAC AAA AGT TGG GAG AAG AAA ATC CAG GAA GCC ATT GAC ATG AGG 4568 Asp Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480 1485 1490  ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC 4613 Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1500 1505  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658 Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510 1515  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG TYR Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG ACG TTG AAA ATC AAC CAG GCA GCA AAC GAA CAA ATA TGC CTA TAC GCG ATA CTG TAC TTG TAC TTG TAC TTG TAC TTG TAC TTG ACG ACG ACG ACG ACG ACG ACG ACG ACG AC
ASP Lys Ser Trp Glu Lys Lys Ile Gln Glu Ala Ile Asp Met Arg 1480  ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1495  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC 4613 Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1500  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658 Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1520  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG 4703 Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 4748 Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793 Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838 Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
ACG GCT GTG GAG TTG CTC AAT GAT GAC GTG GAG CTG ACC ACA GAC 4613 Thr Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr Thr Asp 1500  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG 4703 Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG TTP Ala 1555  CTG GCC GAA ACA ATG GAC AAC AAC GAA CAG ATA TGC CTA TAC GCG 4793 TSSS
The Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr The Asp 1505  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658 Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Thr Asp Gly 1530  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 1530  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 1540  TCG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793 Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
The Ala Val Glu Leu Leu Asn Asp Asp Val Glu Leu Thr The Asp 1505  TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658 Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Thr Asp Gly 1530  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 1530  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 1540  TCG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793 Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658  Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510 1515 1520  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG 4703  Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525 1530 1535  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 4748  Lys Phe Asn Gln Ala Ala 11e Asp Met Ala Glu 11e Leu Thr Leu 1540 1555 1560 1565  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793  TTP Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
TTG GTG AGA GTG CAC CCG GAC AGC AGC CTG GTG GGT CGT AAG GGC 4658  Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510 1515 1520  TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG 4703  Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525 1530 1535  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 4748  Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540 1545 1550  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793  Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510
Leu Val Arg Val His Pro Asp Ser Ser Leu Val Gly Arg Lys Gly 1510
TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
TAC AGT ACC ACT GAC GGG TCG CTG TAC TCG TAC TTT GAA GGT ACG Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525 1530 1535  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540 1545 1550  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838 Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Tyr Ser Thr Thr Asp Gly Ser Leu Tyr Ser Tyr Phe Glu Gly Thr 1525  AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG TTP Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG 4748 Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540 1550  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793 Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838 Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
AAA TTC AAC CAG GCT GCT ATT GAT ATG GCA GAG ATA CTG ACG TTG Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540 1545 1550  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793 Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838 Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793  Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Lys Phe Asn Gln Ala Ala Ile Asp Met Ala Glu Ile Leu Thr Leu 1540  TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793  Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793  Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
TGG CCC AGA CTG CAA GAG GCA AAC GAA CAG ATA TGC CTA TAC GCG 4793  Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838  Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Trp Pro Arg Leu Gln Glu Ala Asn Glu Gln Ile Cys Leu Tyr Ala 1555  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
1555 1560 1565  CTG GGC GAA ACA ATG GAC AAC ATC AGA TCC AAA TGT CCG GTG AAC 4838 Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Leu Gly Glu Thr Met Asp Asn Ile Arg Ser Lys Cys Pro Val Asn 1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
1570 1757 1580  GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
GAT TCC GAT TCA TCA ACA CCT CCC AGG ACA GTG CCC TGC CTG TGC 4883
Asp Ser Asp Ser Ser Thr Pro Pro Arg Thr Val Pro Cys Leu Cys
1585 1590 1595
CGC TAC GCA ATG ACA GCA GAA CGG ATC GCC CGC CTT AGG TCA CAC 4928
Arg Tyr Ala Met Thr Ala Glu Arg Ile Ala Arg Leu Arg Ser His
1600 1605 1610
CAA GTT AAA AGC ATG GTG GTT TGC TCA TCT TTT CCC CTC CCG AAA 4973
Gln Val Lys Ser Met Val Val Cys Ser Ser Phe Pro Leu Pro Lys
1615 1620 1625
THE CAT CTA CAT CCC CTC CAG AAG GTA AAG TGC GAG AAG GTT CTC 5018
TAC CAT GIA GAT GGG GIG CAG ALLO GIA TELO TO
Tyr His Val Asp Gly Val Gln Lys Val Lys Cys Glu Lys Val Leu 1630 1635 1640
1630 1635 1640

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### Figure 5 (9)

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ACG ACG GAC Thr Thr Asp 1665			
TGG ACC ACC Trp Thr Thr 1680			
CCC AGT TTG Pro Ser Leu 1695		•	
GCT CCC ATA Ala Pro Ile 1710			
ATC GCG GAC Ile Ala Asp 1725			
GTG GAC CTC Val Asp Leu 1740			
GCA TAC CTT Ala Tyr Leu 1755			
AGA AAG CCG Arg Lys Pro 1770			
CCT TTG ACG Pro Leu Thr 1785			
 GCC TCC GGG Ala Ser Gly 1800			
GGC CGC GCG Gly Arg Ala 1815			
CAT TTA CAA His Leu Gln 1830			
CAA CTG GAT Gln Leu Asp 1845			
GAT ACT GAG Asp Thr Glu 1860			

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2065

CA Gl	G ATO	G CAG	C CC	A TCC	G GAG	G GCT	T AAT	r AAc	G AG	T CG	A TAC	CAC	G TC	r CGC	5738
	70				1875					188		. 011	ı se	AIG	
														A TCG	
18	85				1890	)				189	5			r Ser	
GG(	G GCC Y Ala	: AGA	A TTO g Leo	G TAC	ACC Thr	GGA Gly	A GCC	G GAC	GT.	A GGC	CGC	ATA	A CCA	A ACA	5828
19	00				1905	•	•			1910	0				
														ATC lle	5873
19				, -,-	1920		,	, vu	y.	1925		, 1111	. val	. IIe	
														GAA	5918
19:		Pne	e Ser	Ser	1935		val	. Ala	ı Ile	≥ Ala 1940		Cys	: Asn	Glu	
TAC	CTA	TCC	AGA	AAT	TAC	CCA	ACA	GTG	GCC	TCG	TAC	CAG	ATA	ACA	5963
191		Ser	. Arg		1950	Pro	Thr	· Val	Ala	1955		Gln	lle	Thr	
														AGT	6008
196	Glu 50	туг	Asp		1yr 1965	reu	Asp	Met	Val	1970		Ser	Asp	Ser	
														TAC	6053
197	Leu 75	Asp	Arg		1nr 1980	Pne	Cys	Pro	Ala	Lys 1985		Arg	Cys	Tyr	
	AAA														6098
199	Lys 0	HIS	HIS		1yr 1995	Hls	GIn	Pro	Thr	Val 2000		Ser	Ala	Val	
	TCA														6143
200	Ser 5	Pro	Pne		Asn 2010	Thr	Leu	GIn	Asn	Val 2015		Ala	Ala	Ala	
	AAG														6188
202	Lys 0	Arg	Asn		Asn 2025	Val	Thr	Gln		Arg 2030	Glu	Leu	Pro	Thr	
	GAC														6233
мес 203	Asp 5	ser	WIG		Phe 2040	ASN	val	Glu		Phe 2045	Lys	Arg	Tyr	Ala	
	TCC														6278
cys 205	Ser 0	GIÀ	GIU		Trp :055	GIu	GIU	Tyr		Lys 2060	Gln	Pro	Ile	Arg	
	ACC														6323
110	Thr	Thr	Glu	Asn	Tle	Thr	Thr	Tyr	Val	Thr	INC	I.a.ı	Tuc	Clv	

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### Figure 5 (11)

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CCG AAA GCT GCT Pro Lys Ala Ala 2080	GCC TTG TTC ( Ala Leu Phe 2 2085	GCT AAG ACC CAC Ala Lys Thr His 2090	AAC TTG GTT CCG 6368 Asn Leu Val Pro
CTG CAG GAG GTT Leu Gln Glu Val 2095	CCC ATG GAC A Pro Met Asp A 2100	AGA TTC ACG GTC Arg Phe Thr Val 2105	GAC ATG AAA CGA 6413 Asp Met Lys Arg
GAT GTC AAA GTC Asp Val Lys Val 2110	ACT CCA GGG 7 Thr Pro Gly 2 2115	ACG AAA CAC ACA Thr Lys His Thr 2120	GAG GAA AGA CCC 6458 Glu Glu Arg Pro
AAA GTC CAG GTA Lys Val Gln Val 2125	ATT CAA GCA ( Ile Gln Ala . 2130	GCG GAG CCA TTG Ala Glu Pro Leu 2135	GCG ACC GCT TAC 6503 Ala Thr Ala Tyr
CTG TGC GGC ATC Leu Cys Gly Ile 2140	His Arg Glu 2145	Leu Val Arg Arg 2150	Leu Asn Ala Val
Leu Arg Pro Asn 2155	Val His Thr 2160	Leu Phe Asp Met 2165	
Phe Asp Ala Ile 2170	Ile Ala Ser 2175	His Phe His Pro 2180	
Leu Glu Thr Asp 2185	Ile Ala Ser 2190	Phe Asp Lys Ser 2195	,
Leu Ala Leu Thr 2200	Gly Leu Met 2205	Ile Leu Glu Asp 2210	CTA GGG GTG GAT 6728 Leu Gly Val Asp
Gln Tyr Leu Leu 2215	Asp Leu Ile 2220	Glu Ala Ala Phe 2225	GGG GAA ATA TCC 6773 Gly Glu Ile Ser
AGC TGT CAC CTA Ser Cys His Leu 2230	CCA ACT GGC Pro Thr Gly 2235	ACG CGC TTC AAG Thr Arg Phe Lys 2240	TTC GGA GCT ATG 6818 Phe Gly Ala Met
ATG AAA TCG GGC Met Lys Ser Gly 2245	ATG TTT CTG Met Phe Leu 2250	ACT TIG TIT ATT Thr Leu Phe Ile 2255	AAC ACT GTT TTG 6863 Asn Thr Val Leu
AAC ATC ACC ATA Asn Ile Thr Ile 2260	GCA AGC AGG Ala Ser Arg 2265	GTA CTG GAG CAG Val Leu Glu Gln 2270	AGA CTC ACT GAC 6908 Arg Leu Thr Asp
TCC GCC TGT GCG Ser Ala Cys Ala 2275	GCC TTC ATC Ala Phe Ile 2280	GGC GAC GAC AAC Gly Asp Asp Asn 2285	ATC GTT CAC GGA 6953 Ile Val His Gly

# . Figure 5 (12)

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GTG ATC TCC Val Ile Ser 2290	C GAC AAG C r Asp Lys L 229	eu Met Ala	GAG AGG TGG Glu Arg Cys 2300	C GCG TCG TGG S Ala Ser Trp	GTC 6998 Val
AAC ATG GAG Asn Met Glu 2305	G GTG AAG A 1 Val Lys I 231	le Ile Asp	GCT GTC ATC Ala Val Met 2315	GGC GAA AAA Gly Glu Lys	CCC 7043 Pro
CCA TAT TTT Pro Tyr Phe 2320	TGT GGG GG Cys Gly Gl 232	y Phe Ile	GTT TTT GAC Val Phe Asp 2330	AGC GTC ACA Ser Val Thr	CAG 7088 Gln
		r Asp Pro		CTG TTC AAG Leu Phe Lys	
GGT AAG CCG Gly Lys Pro 2350	CTA ACA GO Leu Thr Al 235	a Glu Asp	AAG CAG GAC Lys Gln Asp 2360	GAA GAC AGG Glu Asp Arg	CGA 7178 Arg
CGA GCA CTG Arg Ala Leu 2365	AGT GAC GA Ser Asp Gl 237	u Val Ser	AAG TGG TTC Lys Trp Phe 2375	CGG ACA GGC Arg Thr Gly	TTG 7223 Leu
GGG GCC GAA Gly Ala Glu 2380	CTG GAG GT Leu Glu Va 238	l Ala Leu	ACA TCT AGG Thr Ser Arg 2390	TAT GAG GTA Tyr Glu Val	GAG 7268 Glu
GGC TGC AAA Gly Cys Lys 2395	AGT ATC CTO Ser Ile Leo 2400	ı Ile Ala	ATG ACC ACC Met Thr Thr 2405	TTG GCG AGG Leu Ala Arg	GAC 7313 Asp
		Leu Arg		ATA CAC CTC	
GGC GGT CCT Gly Gly Pro 2425	AGA TTG GTG Arg Leu Val 2430	. Arg	TACACAGAAT 1	CTGATTATA GCO	GCACTATT 7412
AȚAGCACC ATG Met	AAT TAC AT Asn Tyr Il	C CCT ACG e Pro Thr 5	CAA ACG TTT Gln Thr Phe	TAC GGC CGC Tyr Gly Arg	CGG 7459 Arg
TGG CGC CCG Trp Arg Pro	CGC CCG GCG Arg Pro Ala	GCC CGT (Ala Arg I	CCT TGG CCG	TTG CAG GCC A Leu Gln Ala T 25	ACT 7504 Thr
CCG GTG GCT Pro Val Ala 30					
CAA CTC ATC AGIn Leu Ile :					

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	GCT Ala 60	Pro							 ACC Thr	7639
	CCA Pro 75	Lys						_	ACG Thr	7684
_	CAG Gln 90						•			7729
_	CCC Pro 105			_			_	_		7774
	ATC Ile 120									7819
	CTG Leu 135									7864
	ATC Ile 150									7909
	AAG Lys 165									7954
	GAT Asp 180									7999
	TGG Trp 195									8044
	CCG Pro 210									8089
	GAC Asp 225	. <i>i</i> .								8134
	GAG Glu 240									8179
GAT Asp	ATG		 							8224





### Figure 5 (14)

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			ATT Ile											TTC Phe	8269
			CAG Gln												8314
			ACA Thr												8359
			GAC Asp											GGA Gly	8404
			CGG Arg					_		_				AAG Lys	8449
			CCT Pro												8494
			CAT His												8539
			GGG Gly												8584
			AGT Ser											GCA Ala	8629
			GCC Ala												8674
			GGA Gly											TTC Phe	8719
			AAG Lys											ATC Ile	8764
			AGA Arg											CAT His	8809
CAT His	CAC	ССТ	CDD	CCG	GTG	CCT	AGA	GAA	AAA	ארעע	ACA	АТТ	AGA	CCA	8854

### Figure 5 (15)

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		AAA Lys						ACA Thr	8899
		GTG Val						ACG Thr	8944
		ACG Thr						ATC Ile	8989
		GGA Gly							9034
		GGC Gly						TGT Cys	9079
		CAG Gln							9124
		TCA Ser							9169
		GTC Val							9214
		ATG Met						AGA Arg	9259
		CTG Leu	Leu			Pro			9304
		CTG Leu						GTG Val	9349
	 	GTG Val						ATG Met	9394
		TGG Trp						CAA Gln	9439
Leu		GAA Glu							9484

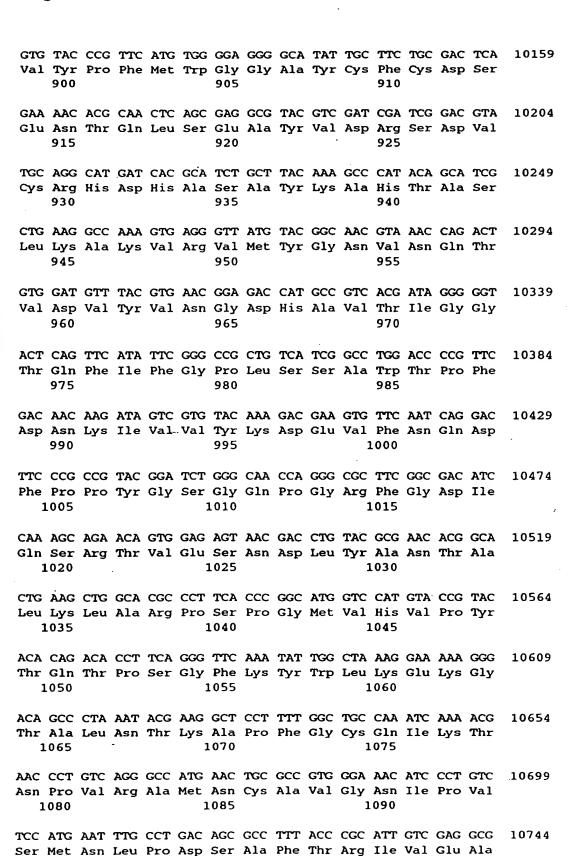
### Figure 5 (16)

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TAC Tyr 690								9529
GGG Gly 705								9574
ATG Met 720								9619
ACA Thr 735								9664
GCC Ala 750					Ala		GCC Ala	9709
TTG Leu 765								9754
CCT Pro 780							AAC Asn	9799
CTG Leu 795								9844
GGG Gly 810								9889
GTG Val 825								9934
AGC Ser 840								9979
CCA Pro 855								10024
GTC Val 870								10069

21/33

1095



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1105

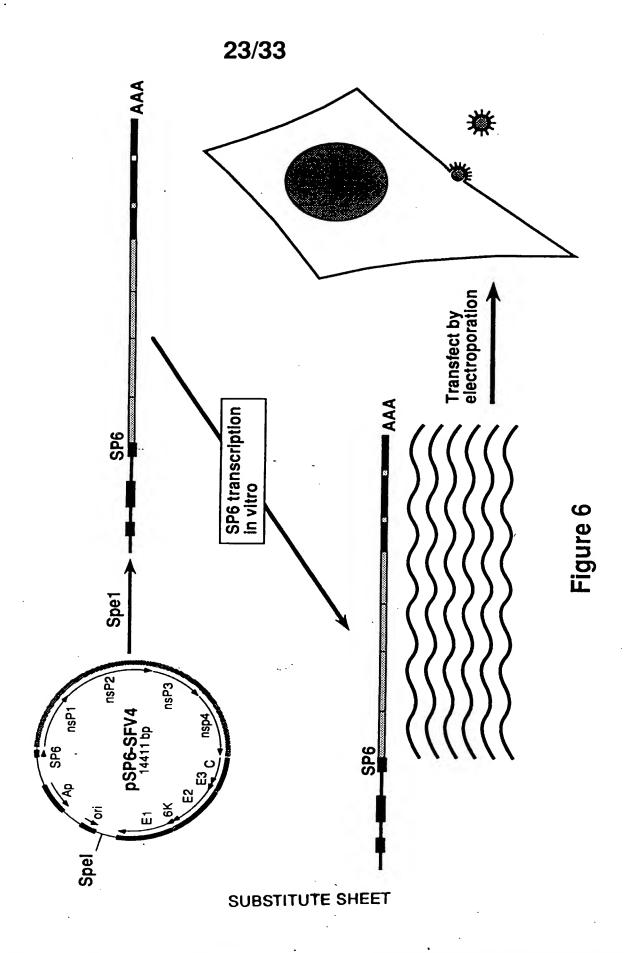
HODEKO. DOFFORE



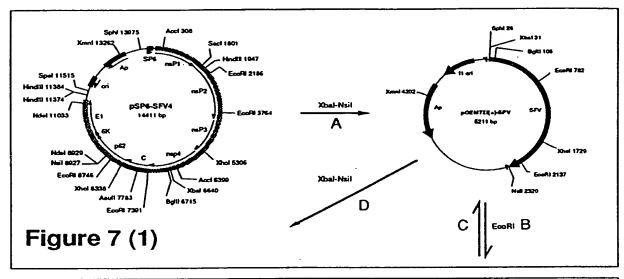
# Figure 5 (18)

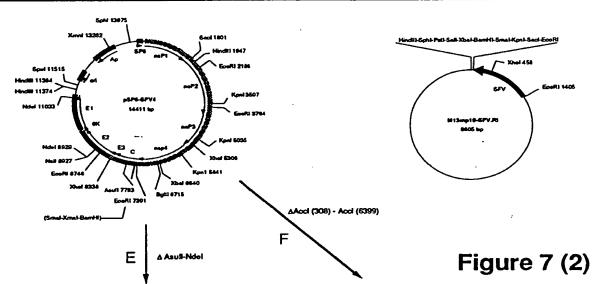
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CCG ACC ATC ATT GAC CTG ACT TGC ACA GTG GCT ACC TGT ACG CAC 10789  Pro Thr Ile Ile Asp Leu Thr Cys Thr Val Ala Thr Cys Thr His  1110 1115 1120	
TCC TCG GAT TTC GGC GGC GTC TTG ACA CTG ACG TAC AAG ACC AAC  Ser Ser Asp Phe Gly Gly Val Leu Thr Leu Thr Tyr Lys Thr Asn  1125  1130  1135	
AAG AAC GGG GAC TGC TCT GTA CAC TCG CAC TCT AAC GTA GCT ACT Lys Asn Gly Asp Cys Ser Val His Ser His Ser Asn Val Ala Thr 1140 1145 1150	
CTA CAG GAG GCC ACA GCA AAA GTG AAG ACA GCA GGT AAG GTG ACC Leu Gln Glu Ala Thr Ala Lys Val Lys Thr Ala Gly Lys Val Thr 1155 1160 1165	
TTA CAC TTC TCC ACG GCA AGC GCA TCA CCT TCT TTT GTG GTG TCG 10969  Leu His Phe Ser Thr Ala Ser Ala Ser Pro Ser Phe Val Val Ser  1170 1175 1180	
CTA TGC AGT GCT AGG GCC ACC TGT TCA GCG TCG TGT GAG CCC CCG 11014  Leu Cys Ser Ala Arg Ala Thr Cys Ser Ala Ser Cys Glu Pro Pro  1185 1190 1195	
AAA GAC CAC ATA GTC CCA TAT GCG GCT AGC CAC AGT AAC GTA GTG Lys Asp His Ile Val. Pro Tyr Ala Ala Ser His Ser Asn Val Val 1200 1205 1210	
TTT CCA GAC ATG TCG GGC ACC GCA CTA TCA TGG GTG CAG AAA ATC  Phe Pro Asp Met Ser Gly Thr Ala Leu Ser Trp Val Gln Lys Ile  1215  1220  1225	
TCG GGT GGT CTG GGG GCC TTC GCA ATC GGC GCT ATC CTG GTG CTG  Ser Gly Gly Leu Gly Ala Phe Ala Ile Gly Ala Ile Leu Val Leu  1230 1235 1240	
GTT GTG GTC ACT TGC ATT GGG CTC CGC AGA TAA GTTAGGGTAG  Val Val Val Thr Cys Ile Gly Leu Arg Arg  1245  1250	
GCAATGGCAT TGATATAGCA AGAAAATTGA AAACAGAAAA AGTTAGGGTA AGCAATGGCA	11252
TATAACCATA ACTGTATAAC TTGTAACAAA GCGCAACAAG ACCTGCGCAA TTGGCCCCGT	11312
GGTCCGCCTC ACGGAAACTC GGGGCAACTC ATATTGACAC ATTAATTGGC AATAATTGGA	11372
AGCTTACATA AGCTTAATTC GACGAATAAT TGGATTTTTA TTTTATTTTG CAATTGGTTT	11432
ТТААТАТТТС САААААААА ААААААААА ААААААААА	11492
АААААААА ААААААААА АСТАG	11517



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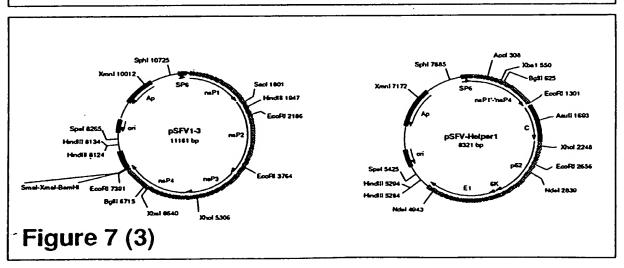


Figure 7 layout scheme

**Figure 7 (1)** 

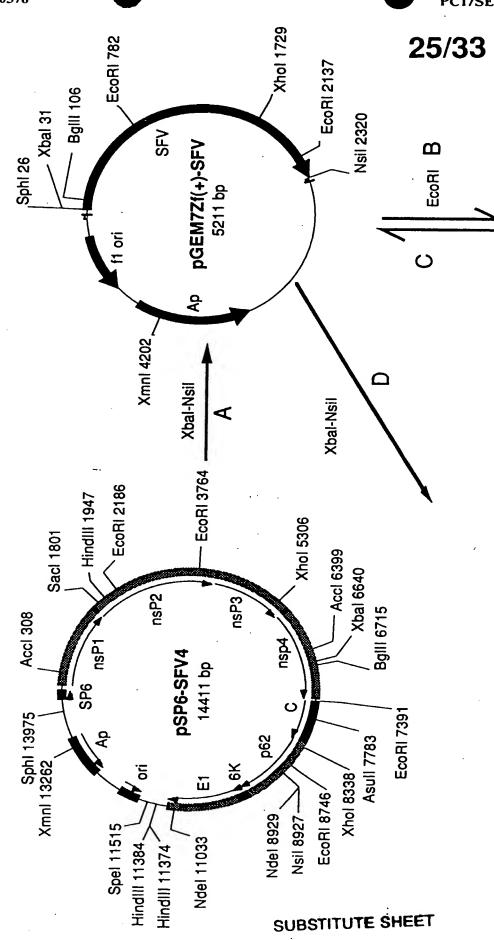
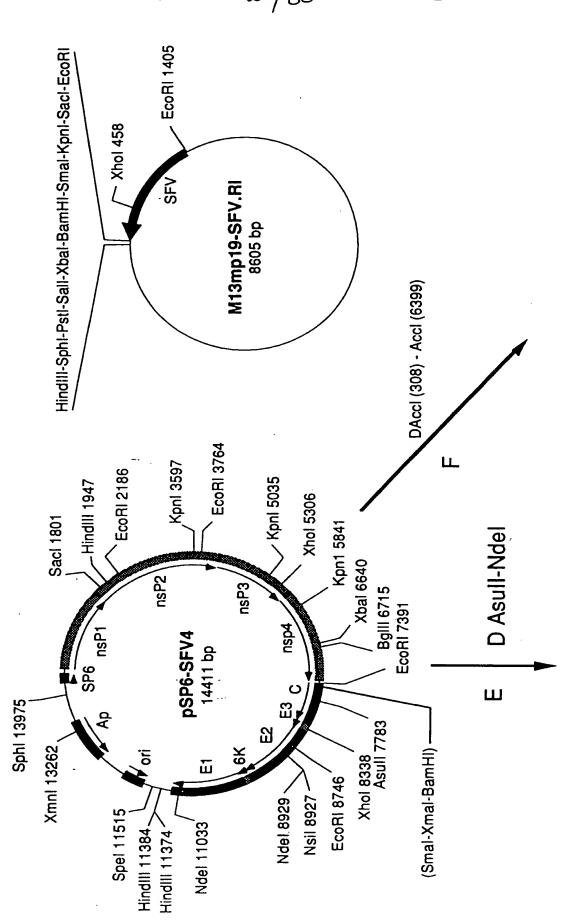
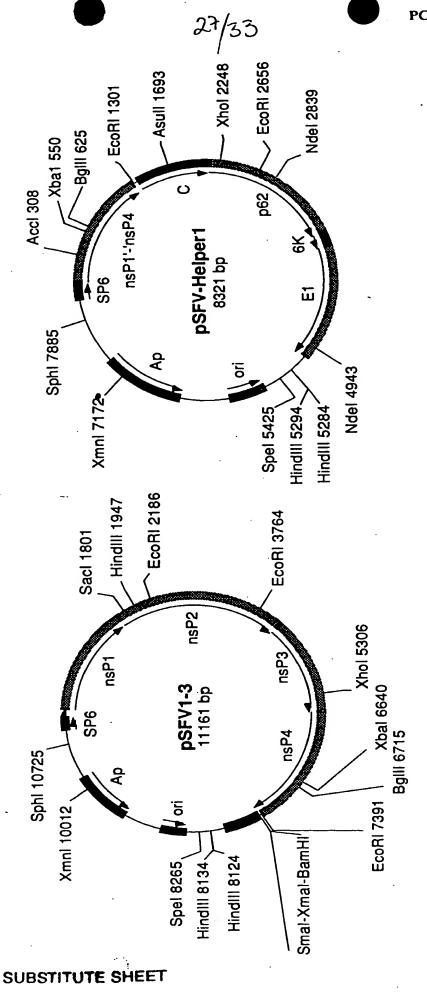


Figure 7 (2)







**Figure 7 (3)** 

SP2

Хтаз

SP1

26S promoter

pSFV2

SP

nsP4

ACCTCTACGG

pSFV3

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Xma3

SP2

26S RNA

26S promoter

pSFV1

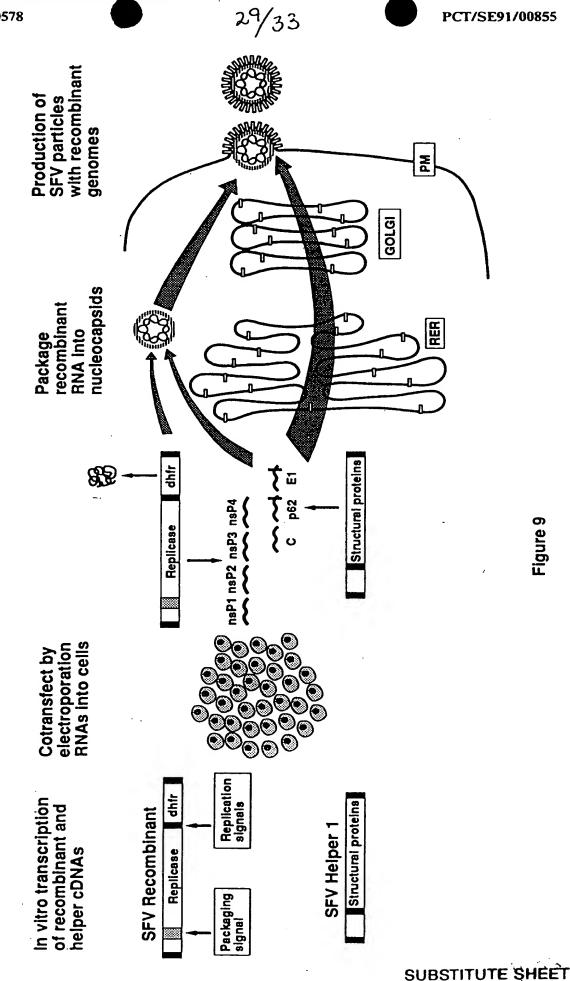
SP1

SP2

Figure 8

DOGGLIOS DZIOGI

Figure 9



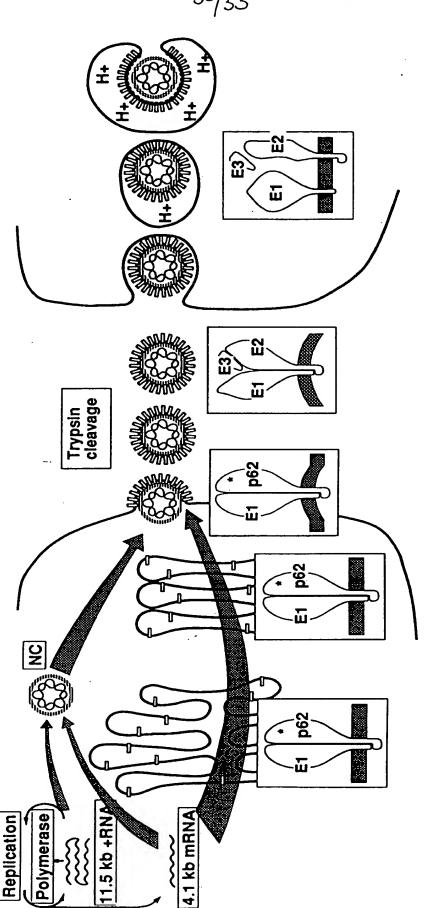


Figure 10

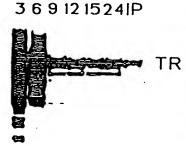
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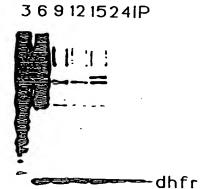


# Receptor

Human Transferrin Mouse Dihydrofolate Reductase

369121524IP -p62





### Chicken Lysozyme

__medlum lysate U 1 2 4 6 8 1 2 4 6 8 IP ∍-lysozyme Transferrin receptor infection/pSFVI

5 .5 .05 .005

Figure 11

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		<del></del>	
AAA GGC AAA GTC CAT SFV vector E2 Lys Gly Lys Val His SFV E2 SFV vector E2	GTT GAG CAA CTC CTA G HIV-gp120 Val Glu Asp HIV epitope	mH1 oligo	GTT GAG GAT CCG CAA CTC CTA GGC Val Glu Asp Pro SFV-HIV chimera
BamH1  AAC TCA CCT TTC GTC CCG AGA GCC GAC GAA CCG GCT AGA A  Asn Ser Pro Phe Val Pro Arg Ala Asp Glu Pro Ala Arg I  Glu Asp	GAT CCG CGT ATC CAG AGA GGA CCA GGA AGA GCA TTT GC GCA TAG GTC TCT CCT GGT CCT TCT CGT AAA ASP Pro Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe	Cut with BamH1 Insert HIV oligo	GAE GAT CCG CGT ATC CAG AGA GGA CCA GGA AGA GCA TTT G CTC CTA GGC GCA TAG GTC TCT CCT GGT CCT TCT CGT AAA C Glu Asp Pro Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe V 313

Figure 12 (1)

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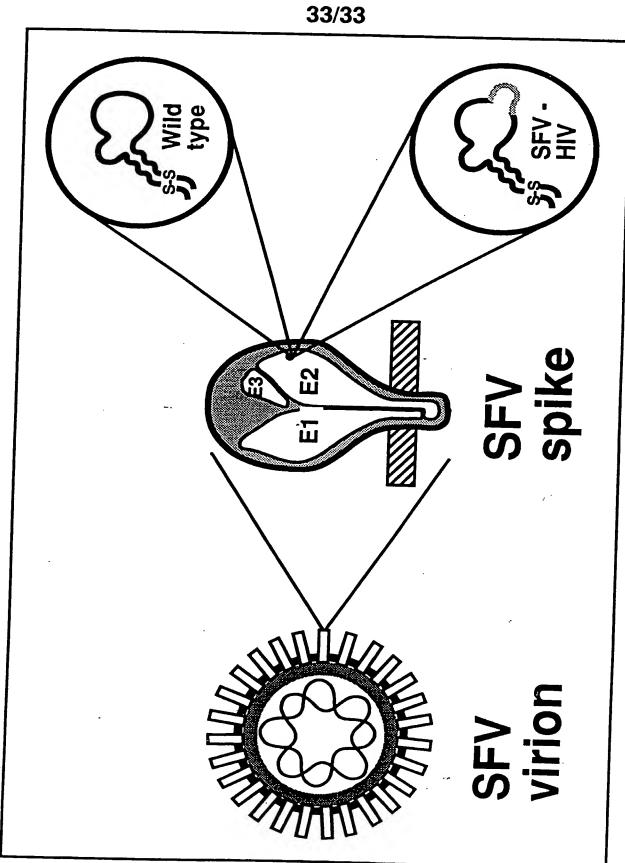


Figure 12 (2)